

1634

#72
KAY
9-17-02

TECH CENTER 1600/2000

AUG 13 2002

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600

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/516,444A

DATE: 08/08/2002

TIME: 12:03:45

Input Set : A:\5995USP2.txt

Output Set: N:\CRF4\08082002\I516444A.raw

```

4 <110> APPLICANT: Abbott Laboratories
5     Billing-Medel, Patricia A.
6     Cohen, Maurice
7     Colpitts, Tracey L.
8     Friedman, Paula N.
9     Gordon, Julian
10    Granados, Edward N.
11    Hodges, Steven C.
12    Klass, Michael R.
13    Kratochvil, Jon D.
14    Roberts-Rapp, Lisa
15    Russell, John C.
16    Stroupe, Stephen D.
18 <120> TITLE OF INVENTION: Reagents And Method Useful For Detecting
19 Diseases Of The Breast
22 <130> FILE REFERENCE: 5995.US.P2
24 <140> CURRENT APPLICATION NUMBER: 09/516,444A
25 <141> CURRENT FILING DATE: 2000-02-29
27 <150> PRIOR APPLICATION NUMBER: US 08/962,094
28 <151> PRIOR FILING DATE: 1997-10-31
30 <150> PRIOR APPLICATION NUMBER: US 08/742,067
31 <151> PRIOR FILING DATE: 1996-10-31
33 <160> NUMBER OF SEQ ID NOS: 39
35 <170> SOFTWARE: FastSEQ for Windows Version 4.0
37 <210> SEQ ID NO: 1
38 <211> LENGTH: 201
39 <212> TYPE: DNA
40 <213> ORGANISM: Artificial Sequence
42 <220> FEATURE:
43 <223> OTHER INFORMATION: EST Clone 1662885
W--> 45 <221> NAME/KEY: misc_feature
46 <222> LOCATION: (26)...(26)
47 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
48 position 26
W--> 50 <221> misc_feature
51 <222> LOCATION: (98)...(98)
52 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
53 position 98
W--> 55 <221> misc_feature
56 <222> LOCATION: (133)...(133)
57 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
58 position 133
W--> 60 <221> misc_feature

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Input Set : A:\5995USP2.txt

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61 <222> LOCATION: (145)...(145)
 62 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
 position 145
 W--> 65 <221> misc_feature
 66 <222> LOCATION: (183)...(183)
 67 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
 position 183
 W--> 70 <400> 1
 W--> 71 ctcttaggct ttgaagcatt tttgtntgtt ctccctgate ttcatgtcac caccatgaag 60
 W--> 72 ttcttagcag tcctggtaact cttggggatt tccatctntc tggctctgc ccagaatccg 120
 W--> 73 acaacagctg ctncagctga cacgnatcca gctactggtc ctgctgatga tgaagcccc 180
 W--> 74 gangctgaaa ccactgctgc t 201
 76 <210> SEQ ID NO: 2
 77 <211> LENGTH: 308
 78 <212> TYPE: DNA
 79 <213> ORGANISM: Homo sapiens
 81 <400> SEQUENCE: 2
 82 taggcttgaa agcatttttg tctgtgctcc ctgatcttca ggtcaccacc atgaagttct 60
 83 tagcagtctt ggtactcttg ggagtttcoa tctttctggg ctctgcccag aatccgacaa 120
 84 cagctgctcc agctgacacg tatccagcta ctggctctgc tgatgatgaa gccctgtatg 180
 85 ctgaaaccac tgctgctgca accactgcga ccactgctgc tcctaccact gcaaccaccg 240
 86 ctgcttctac cactgctcgta aaagacatttcc cagtttacc caaatgggtt gggatcttc 300
 308
 87 cgaatggt
 89 <210> SEQ ID NO: 3
 90 <211> LENGTH: 292
 91 <212> TYPE: DNA
 92 <213> ORGANISM: Artificial Sequence
 94 <220> FEATURE:
 95 <223> OTHER INFORMATION: EST Clone 901429
 W--> 97 <221> NAME/KEY: misc_feature
 98 <222> LOCATION: (236)...(236)
 99 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
 position 236
 W--> 102 <221> misc_feature
 103 <222> LOCATION: (259)...(259)
 104 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
 position 259
 W--> 107 <400> 3
 108 gcatttttgt ctgtgctccc tgatcttcat gtcaccacca tgaagtttctt agcagtccgt 60
 109 gtactcttgg gagtttccat ctttctggc tctgcccaga atccgacaa ac agctgctcca 120
 110 gctgacacgt atccagctac tggctctgtt gatgatgaa cccctgtatgc taaaaccact 180
 W--> 111 gctgctgcaa ccactgctgac cactgctgtt cctaccactg caaccaccgc tgcttntacc 240
 W--> 112 actgctcgta aagacatttnc agtttaccacca aatgggtt gggatctccc ga 292
 114 <210> SEQ ID NO: 4
 115 <211> LENGTH: 197
 116 <212> TYPE: DNA
 117 <213> ORGANISM: Homo sapiens
 119 <400> SEQUENCE: 4
 120 gttttaccca aatgggttgg ggatctcccg aatggtagag tggccctgtt agatggaatc 60

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```

121 agcttgagtc ttctgcaatt ggtcacaact attcatgctt cctgtgatTT catccaacta 120
122 cttaccttgc ctacgatatac ccctttatct ctaatcagtt tattttcttt caaataaaaa 180
123 ataactatga gcaacat 197
125 <210> SEQ ID NO: 5
126 <211> LENGTH: 472
127 <212> TYPE: DNA
128 <213> ORGANISM: Homo sapiens
130 <400> SEQUENCE: 5
131 ctcttaggct ttgaagcatt tttgtctgtg ctccctgatC ttcatgtcac caccatgaag 60
132 ttcttagcag tcctggtaCT cttgggaggT tccatcttC tggtctctgc ccagaatccg 120
133 acaacagctg ctccagctga cacgtatcca gctactggTC ctgctgatga tgaagcccc 180
134 gatgctgaaa ccactgctgc tgcaaccact gcgaccactg ctgctcctac cactgcaacc 240
135 accgctgctt ctaccactgc tcgtaaagac attccagtt tacccaaatg ggttggggat 300
136 ctcccgaatg gtagagtgtg tccctgagat ggaatcagct tgagtcttct gcaattggtc 360
137 acaactattc atgcttcctg tgatttcatc caactactta ccttgcctac gatatcccc 420
138 ttatctctaa tcagtttatt ttctttcaaa taaaaaataa ctatgagcaa ca 472
140 <210> SEQ ID NO: 6
141 <211> LENGTH: 473
142 <212> TYPE: DNA
143 <213> ORGANISM: Homo sapiens
145 <400> SEQUENCE: 6
146 ctcttaggct ttgaagcatt tttgtctgtg ctccctgatC ttcatgtcac caccatgaag 60
147 ttcttagcag tcctggtaCT cttgggaggT tccatcttC tggtctctgc ccagaatccg 120
148 acaacagctg ctccagctga cacgtatcca gctactggTC ctgctgatga tgaagcccc 180
149 gatgctgaaa ccactgctgc tgcaaccact gcgaccactg ctgctcctac cactgcaacc 240
150 accgctgctt ctaccactgc tcgtaaagac attccagtt tacccaaatg ggttggggat 300
151 ctcccgaatg gtagagtgtg tccctgagat ggaatcagct tgagtcttct gcaattggtc 360
152 acaactattc atgcttcctg tgatttcatc caactactta ccttgcctac gatatcccc 420
153 ttatctctaa tcagtttatt ttctttcaaa taaaaaataa ctatgagcaa cat 473
155 <210> SEQ ID NO: 7
156 <211> LENGTH: 68
157 <212> TYPE: DNA
158 <213> ORGANISM: Artificial Sequence
160 <220> FEATURE:
161 <223> OTHER INFORMATION: Restriction site
163 <400> SEQUENCE: 7
164 agctcggaat tccgagcttG gatcctctAG agcggccGCC gactagtGAG ctcgtcgacc 60
165 cgggatt 68
167 <210> SEQ ID NO: 8
168 <211> LENGTH: 68
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
172 <220> FEATURE:
173 <223> OTHER INFORMATION: Restriction site
175 <400> SEQUENCE: 8
176 aattaattcc cgggtcgacg agctcaCTAG tcggcggCCG ctctagagGA tccaaGCTG 60
177 gaattccG 68
179 <210> SEQ ID NO: 9
180 <211> LENGTH: 24

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RAW SEQUENCE LISTING

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Input Set : A:\5995USP2.txt
 Output Set: N:\CRF4\08082002\I516444A.raw

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181 <212> TYPE: DNA
182 <213> ORGANISM: Artificial Sequence
184 <220> FEATURE:
185 <223> OTHER INFORMATION: Universal Primer
187 <400> SEQUENCE: 9
188 agcggataac aatttcacac agga 24
190 <210> SEQ ID NO: 10
191 <211> LENGTH: 18
192 <212> TYPE: DNA
193 <213> ORGANISM: Artificial Sequence
195 <220> FEATURE:
196 <223> OTHER INFORMATION: Universal Primer
198 <400> SEQUENCE: 10
199 tgtaaaacgca cggccagt 18
201 <210> SEQ ID NO: 11
202 <211> LENGTH: 20
203 <212> TYPE: DNA
204 <213> ORGANISM: Artificial Sequence
206 <220> FEATURE:
207 <223> OTHER INFORMATION: Primer
209 <400> SEQUENCE: 11
210 actgctcgta aagacattcc 20
212 <210> SEQ ID NO: 12
213 <211> LENGTH: 19
214 <212> TYPE: DNA
215 <213> ORGANISM: Artificial Sequence
217 <220> FEATURE:
218 <223> OTHER INFORMATION: Primer
220 <400> SEQUENCE: 12
221 gggacacact ctaccattc 19
223 <210> SEQ ID NO: 13
224 <211> LENGTH: 20
225 <212> TYPE: DNA
226 <213> ORGANISM: Artificial Sequence
228 <220> FEATURE:
229 <223> OTHER INFORMATION: Sense Primer
231 <400> SEQUENCE: 13
232 aagccccctgat tgctgaaacc 20
234 <210> SEQ ID NO: 14
235 <211> LENGTH: 23
236 <212> TYPE: DNA
237 <213> ORGANISM: Artificial Sequence
239 <220> FEATURE:
240 <223> OTHER INFORMATION: Antisense Primer
242 <400> SEQUENCE: 14
243 tgcagaagac tcaagctgtat tcc 23
245 <210> SEQ ID NO: 15
246 <211> LENGTH: 20
247 <212> TYPE: DNA

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RAW SEQUENCE LISTING
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Input Set : A:\5995USP2.txt
Output Set: N:\CRF4\08082002\I516444A.raw

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248 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: Target-Specific Forward Primer
253 <400> SEQUENCE: 15
254 aaggccctga tgctgaaacc 20
256 <210> SEQ ID NO: 16
257 <211> LENGTH: 23
258 <212> TYPE: DNA
259 <213> ORGANISM: Artificial Sequence
261 <220> FEATURE:
262 <223> OTHER INFORMATION: Target-Specific Reverse Primer
264 <400> SEQUENCE: 16
265 tgcagaagac tcaagctgat tcc 23
267 <210> SEQ ID NO: 17
268 <211> LENGTH: 15
269 <212> TYPE: DNA
270 <213> ORGANISM: Artificial Sequence
272 <220> FEATURE:
273 <223> OTHER INFORMATION: Probe
275 <400> SEQUENCE: 17
276 gaccactgct gctcc 15
278 <210> SEQ ID NO: 18
279 <211> LENGTH: 20
280 <212> TYPE: DNA
281 <213> ORGANISM: Artificial Sequence
283 <220> FEATURE:
284 <223> OTHER INFORMATION: Sense Primer
286 <400> SEQUENCE: 18
287 actgctcgta aagacattcc 20
289 <210> SEQ ID NO: 19
290 <211> LENGTH: 19
291 <212> TYPE: DNA
292 <213> ORGANISM: Artificial Sequence
294 <220> FEATURE:
295 <223> OTHER INFORMATION: Antisense Primer
297 <400> SEQUENCE: 19
298 gggacacact ctaccattc 19
300 <210> SEQ ID NO: 20
301 <211> LENGTH: 90
302 <212> TYPE: PRT
303 <213> ORGANISM: Homo sapiens
305 <400> SEQUENCE: 20
306 Met Lys Phe Leu Ala Val Leu Val Leu Gly Val Ser Ile Phe Leu
307 1 5 10 15
308 Val Ser Ala Gln Asn Pro Thr Thr Ala Ala Pro Ala Asp Thr Tyr Pro
309 20 25 30
310 Ala Thr Gly Pro Ala Asp Asp Glu Ala Pro Asp Ala Glu Thr Thr Ala
311 35 40 45
312 Ala Ala Thr Thr Ala Thr Ala Ala Pro Thr Thr Ala Thr Thr Ala

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VERIFICATION SUMMARY
PATENT APPLICATION: US/09/516,444A

DATE: 08/08/2002
TIME: 12:03:46

Input Set : A:\5995USP2.txt
Output Set: N:\CRF4\08082002\I516444A.raw

L:45 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:50 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:60 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:65 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:70 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:71 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:60
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:120
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180
L:97 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:102 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:107 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:180
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:240



Creation date: 11-14-2003

Indexing Officer: QNGUYEN5 - QUANG NGUYEN

Team: OIPEBackFileIndexing

Dossier: 09516444

Legal Date: 10-02-2003

No.	Doccode	Number of pages
1	CTRS	6
2	NFDR	1
3	EXIN	1

Total number of pages: 8

Remarks:

Order of re-scan issued on